

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/787,219A
Source: IFWO
Date Processed by STIC: 10/29/04

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IFWO

RAW SEQUENCE LISTING

DATE: 10/29/2004

PATENT APPLICATION: US/10/787,219A

TIME: 12:42:39

Input Set : A:\248628US0X.txt

Output Set: N:\CRF4\10292004\J787219A.raw

3 <110> APPLICANT: JESTIN, JEAN-LUC
 4 VICHIER-GUERRE, SOPHIE
 6 <120> TITLE OF INVENTION: METHODS FOR OBTAINING THERMOSTABLE ENZYMES, DNA POLYMERASE I
 7 VARIANTS FROM THERMUS AQUATICUS HAVING NEW CATALYTIC ACTIVITIES,
 8 METHODS FOR OBTAINING THE SAME, AND APPLICATIONS OF THE SAME
 10 <130> FILE REFERENCE: 248628USOX
 12 <140> CURRENT APPLICATION NUMBER: 10/787,219A
 13 <141> CURRENT FILING DATE: 2004-02-27
 15 <160> NUMBER OF SEQ ID NOS: 61
 17 <170> SOFTWARE: PatentIn version 3.3
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 24
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Synthetic DNA
 27 <400> SEQUENCE: 1
 28 taacaatagg ccggccaccc cttc 24
 31 <210> SEQ ID NO: 2
 32 <211> LENGTH: 18
 33 <212> TYPE: DNA
 34 <213> ORGANISM: Artificial Sequence
 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: Synthetic DNA
 39 <400> SEQUENCE: 2
 40 gagtttttgt tctgcggc 18
 43 <210> SEQ ID NO: 3
 44 <211> LENGTH: 27
 45 <212> TYPE: DNA
 46 <213> ORGANISM: Artificial Sequence
 48 <220> FEATURE:
 49 <223> OTHER INFORMATION: Synthetic DNA
 51 <400> SEQUENCE: 3
 52 tttaatcatc tgcagtaccg ggagctc 27
 55 <210> SEQ ID NO: 4
 56 <211> LENGTH: 28
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Artificial Sequence
 60 <220> FEATURE:
 61 <223> OTHER INFORMATION: Synthetic DNA
 63 <400> SEQUENCE: 4
 64 ttcattcttg ctagctcctg ggagaggc 28
 67 <210> SEQ ID NO: 5

(ps.10)

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68 <211> LENGTH: 43
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Synthetic DNA
76 <220> FEATURE:
77 <221> NAME/KEY: misc_feature
78 <222> LOCATION: (15)..(15)
79 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
80      C and A, in the trimer sequence CAR and AVY, respectively
82 <220> FEATURE:
83 <221> NAME/KEY: misc_feature
84 <222> LOCATION: (16)..(16)
85 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
86      A and V, in the trimer sequence CAR and AVY, respectively
88 <220> FEATURE:
89 <221> NAME/KEY: misc_feature
90 <222> LOCATION: (17)..(17)
91 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
92      R and Y, in the trimer sequence CAR and AVY, respectively
94 <220> FEATURE:
95 <221> NAME/KEY: misc_feature
96 <222> LOCATION: (24)..(24)
97 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
98      C and A, in the trimer sequence CAR and AVY, respectively
100 <220> FEATURE:
101 <221> NAME/KEY: misc_feature
102 <222> LOCATION: (25)..(25)
103 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
104      A and V, in the trimer sequence CAR and AVY, respectively
106 <220> FEATURE:
107 <221> NAME/KEY: misc_feature
108 <222> LOCATION: (26)..(26)
109 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
110      R and Y, in the trimer sequence CAR and AVY, respectively
112 <400> SEQUENCE: 5
W--> 113 ccggccaccc cttcnnnctc aacnnncggg accagctgga aag      43
116 <210> SEQ ID NO: 6
117 <211> LENGTH: 65
118 <212> TYPE: DNA
119 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Synthetic DNA
125 <220> FEATURE:
126 <221> NAME/KEY: misc_feature
127 <222> LOCATION: (17)..(17)
128 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative

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abundance:

129 Y and R, in the trimer sequence YTG and RBT, respectively

131 <220> FEATURE:

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Input Set : A:\248628US0X.txt

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132 <221> NAME/KEY: misc_feature
 133 <222> LOCATION: (18)..(18)
 134 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 135 T and B, in the trimer sequence YTG and RBT, respectively
 137 <220> FEATURE:
 138 <221> NAME/KEY: misc_feature
 139 <222> LOCATION: (19)..(19)
 140 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 141 G and T, in the trimer sequence YTG and RBT, respectively
 143 <220> FEATURE:
 144 <221> NAME/KEY: misc_feature
 145 <222> LOCATION: (20)..(20)
 146 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 147 Y and R, in the trimer sequence YTG and RBT, respectively
 149 <220> FEATURE:
 150 <221> NAME/KEY: misc_feature
 151 <222> LOCATION: (21)..(21)
 152 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 153 T and B, in the trimer sequence YTG and RBT, respectively
 155 <220> FEATURE:
 156 <221> NAME/KEY: misc_feature
 157 <222> LOCATION: (22)..(22)
 158 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 159 G and T, in the trimer sequence YTG and RBT, respectively
 161 <220> FEATURE:
 162 <221> NAME/KEY: misc_feature
 163 <222> LOCATION: (26)..(26)
 164 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 165 Y and R, in the trimer sequence YTG and RBT, respectively
 167 <220> FEATURE:
 168 <221> NAME/KEY: misc_feature
 169 <222> LOCATION: (27)..(27)
 170 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 171 T and B, in the trimer sequence YTG and RBT, respectively
 173 <220> FEATURE:
 174 <221> NAME/KEY: misc_feature
 175 <222> LOCATION: (28)..(28)
 176 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 177 G and T, in the trimer sequence YTG and RBT, respectively
 179 <220> FEATURE:
 180 <221> NAME/KEY: misc_feature
 181 <222> LOCATION: (44)..(44)
 182 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 183 Y and R, in the trimer sequence YTG and RBT, respectively

185 <220> FEATURE:
186 <221> NAME/KEY: misc_feature
187 <222> LOCATION: (45)..(45)
188 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
189 T and B, in the trimer sequence YTG and RBT, respectively

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191 <220> FEATURE:
192 <221> NAME/KEY: misc_feature
193 <222> LOCATION: (46)..(46)
194 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
195      G and T, in the trimer sequence YTG and RBT, respectively
197 <400> SEQUENCE: 6
W--> 198 ggatgaggtc cggcaannnn nnaatnnngg tgctcttcag cttnnngagc tcccgtact      60
200 gcagg                                           65
203 <210> SEQ ID NO: 7
204 <211> LENGTH: 62
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: Synthetic DNA
212 <220> FEATURE:
213 <221> NAME/KEY: misc_feature
214 <222> LOCATION: (17)..(17)
215 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
216      C and A, in the trimer sequence CAR and AVY, respectively
218 <220> FEATURE:
219 <221> NAME/KEY: misc_feature
220 <222> LOCATION: (18)..(18)
221 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
222      A and V, in the trimer sequence CAR and AVY, respectively
224 <220> FEATURE:
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: (19)..(19)
227 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
228      R and Y, in the trimer sequence CAR and AVY, respectively
230 <220> FEATURE:
231 <221> NAME/KEY: misc_feature
232 <222> LOCATION: (32)..(32)
233 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
234      C and A, in the trimer sequence CAR and AVY, respectively
236 <220> FEATURE:
237 <221> NAME/KEY: misc_feature
238 <222> LOCATION: (33)..(33)
239 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
240      A and V, in the trimer sequence CAR and AVY, respectively
242 <220> FEATURE:
243 <221> NAME/KEY: misc_feature
244 <222> LOCATION: (34)..(34)
245 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
246      R and Y, in the trimer sequence CAR and AVY, respectively
248 <220> FEATURE:
249 <221> NAME/KEY: misc_feature

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250 <222> LOCATION: (41)..(41)

251 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

252 C and A, in the trimer sequence CAR and AVY, respectively

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254 <220> FEATURE:
 255 <221> NAME/KEY: misc_feature
 256 <222> LOCATION: (42)..(42)
 257 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 258 A and V, in the trimer sequence CAR and AVY, respectively
 260 <220> FEATURE:
 261 <221> NAME/KEY: misc_feature
 262 <222> LOCATION: (43)..(43)
 263 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 264 R and Y, in the trimer sequence CAR and AVY, respectively
 266 <400> SEQUENCE: 7
 W--> 267 caaccagacg gccacgnnna cgggcaggct annnagctcc nnncccaacc tccagaacat 60
 269 cc 62
 272 <210> SEQ ID NO: 8
 273 <211> LENGTH: 43
 274 <212> TYPE: DNA
 275 <213> ORGANISM: Artificial Sequence
 277 <220> FEATURE:
 278 <223> OTHER INFORMATION: Synthetic DNA
 281 <220> FEATURE:
 282 <221> NAME/KEY: misc_feature
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 284 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 285 Y and R, in the trimer sequence YTG and RBT, respectively
 287 <220> FEATURE:
 288 <221> NAME/KEY: misc_feature
 289 <222> LOCATION: (15)..(15)
 290 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 291 T and B, in the trimer sequence YTG and RBT, respectively
 293 <220> FEATURE:
 294 <221> NAME/KEY: misc_feature
 295 <222> LOCATION: (16)..(16)
 296 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 297 G and T, in the trimer sequence YTG and RBT, respectively
 299 <220> FEATURE:
 300 <221> NAME/KEY: misc_feature
 301 <222> LOCATION: (23)..(23)
 302 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 303 Y and R, in the trimer sequence YTG and RBT, respectively
 305 <220> FEATURE:
 306 <221> NAME/KEY: misc_feature
 307 <222> LOCATION: (24)..(24)
 308 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:
 309 T and B, in the trimer sequence YTG and RBT, respectively
 311 <220> FEATURE:
 312 <221> NAME/KEY: misc_feature

313 <222> LOCATION: (25)..(25)

314 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

315 G and T, in the trimer sequence YTG and RBT, respectively

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/787,219A

DATE: 10/29/2004
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Input Set : A:\248628US0X.txt
Output Set: N:\CRF4\10292004\J787219A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 15,16,17,24,25,26
Seq#:6; N Pos. 17,18,19,20,21,22,28,27,28,44,45,46
Seq#:7; N Pos. 17,18,19,22,33,34,41,42,43
Seq#:8; N Pos. 14,15,16,23,24,25
Seq#:9; N Pos. 20,21,22,38,39,40,44,45,46,47,48,49
Seq#:10; N Pos. 20,21,22,29,30,31,44,45,46
Seq#:11; N Pos. 19,20,21,28,29,30

VERIFICATION SUMMARY

DATE: 10/29/2004

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Input Set : A:\248628US0X.txt

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L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:318 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:528 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0